



NAVAL BIODYNAMICS LABORATORY  
NBDL-90R002

**X-RAY ANTHROPOMETRY TRANSFORMATION PROGRAM  
FOR THE HEWLETT-PACKARD 9000/835 COMPUTER**

DOROTHY FRANCIS

Software Documentation

May 1991

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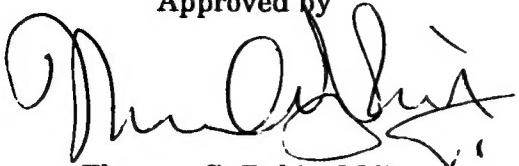
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## TABLE OF CONTENTS

ACKNOWLEDGEMENT .....	ii
1. INTRODUCTION .....	1
2. FUNCTION .....	3
3. MAIN PROGRAM .....	3
3.1 MXRAY .....	3
3.2 COMPILE .....	4
3.3 DATA FILES .....	4
3.4 EXECUTION .....	4
4. SUBROUTINE DESCRIPTIONS .....	5
4.1 AXEB .....	5
4.2 CDNTPT .....	5
4.3 POBKG .....	5
4.4 PXRY2 .....	6
4.5 RLSWCH .....	6
4.6 OPTION .....	7
4.7 PRISM2 .....	7
4.8 HTRANS .....	8
5. UTILITY SUBROUTINES .....	8
5.1 CAPS .....	8
5.2 CAPOFF .....	8
REFERENCE .....	9
APPENDIX A: Program Listings .....	A-1
APPENDIX B: Listing of "IOUPDATE" .....	B-1
APPENDIX C: Listing of "INPUT" .....	C-1
APPENDIX D: Listing of "OUTPUT" .....	D-1
APPENDIX E: Listing of "UPDATE" .....	E-1

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# **X-RAY ANTHROPOMETRY TRANSFORMATION PROGRAM**

## **FOR THE HEWLETT-PACKARD 9000/550 COMPUTER**

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### **1. INTRODUCTION**

The Naval Biodynamics Laboratory (NAVBIODYNLAB), located in New Orleans, Louisiana, is a research facility under the cognizance of the Naval Medical Research and Development Command. It is the only Navy laboratory conducting biomedical research on the effects of mechanical forces (motion and impact) encountered by Navy personnel. Among its goals are the establishment of human tolerance limits and the development of appropriate methods of avoiding and treating the deleterious effects of such forces. Ongoing research programs at the laboratory acquire sensor and photographic impact acceleration data from acceleration sled runs.

To analyze this data, certain anthropometric information about each subject is necessary: the locations of the head and the first thoracic vertebral body (T-1) instrumentation origins relative to the corresponding anatomical origins, as well as transformation matrices from the instrument to the anatomical coordinate systems.

A stereoradiographic technique, in which two simultaneous X-ray exposures of an instrumented subject are prepared and analyzed, is used to measure the geometrical relationships between subject mounted-instrumentation and subject anatomy. This yields a complete six-parameter statement of the position and orientation of the instrument package relative to a coordinate system fixed in the subject's bony anatomy. In the anatomical coordinate system, the mounted instrumentation is assumed to be rigidly fixed. If the geometrical relationship of these systems is known, then the transformation from one system to the other is a matter of rigid body mechanics.

Stereoradiographic techniques have been devised at the Naval Biodynamics Laboratory to measure the geometrical relationships between subject-mounted instrumentation and subject anatomy. Measurements for instrumentation mounts are made at the head and T-1.

Packages consisting of photographic targets and accelerometers that are fixed to a rigid T-shaped plate are attached to human and human analog subjects by means of an intermediary anatomical mount. These mounts are prepared so that one surface provides a rigid and highly repeatable mooring for the instrumentation packages, while another is custom molded to fit the subject's dental surfaces or bony protuberances of the spine or pelvis. The transformation from anatomical to instrumentation coordinates is governed by these anatomical mounts, with a single transformation uniquely associated with each.

Quantization of these transformations is performed radiographically. The subject is X-rayed while wearing lead markers on externally accessible anatomical features. Also worn are anatomical mounts equipped with special Plexiglass™ T-plates, in which additional lead markers ("BBs") have been fixed at known locations. The X-ray images of these markers and of other anatomical features establish the laboratory reference position of the test subject and the orientation of the instrumentation and anatomical coordinate systems. This information is used to convert anatomical to instrumentation data.

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

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Two groups of anatomical coordinates are presently used in the work conducted at NAVBIODYNLAB. One is located in the bony anatomy of the head, the other in the first thoracic vertebral body (T-1). The head coordinate system is defined by the positions of the two auditory meatus and the notches on the inferior orbital ridges. Its origin is the midpoint of the line joining the auditory meatus. The X axis is positive from the origin through the midpoint of the line connecting the orbital notches. The X-Y plane contains the X axis and the midpoints of the lines from the left orbital notch to the left auditory meatus and from the right orbital notch to the right auditory meatus. The Z axis is positive through the top of the skull. The T-1 coordinate system is also based on four points: the posterior tip of the posterior spinous process, the anterior-superior corner of the vertebral body, and the superior corners of the right and left articular facets. Unfortunately, most of these points are not directly accessible for marking and are visible only in certain X-ray projections.

Therefore, the suprasternal notch, an externally accessible site, is used to locate a system with the anterior-superior corner at T-1, with the X axis going through the posterior tip of the posterior spinous process and positive anteriorly, and with the Y axis parallel to the line connecting the two articular facets and positive left. The Z axis is positive up.

Since the radiologic equipment used at NAVBIODYNLAB is primarily intended for diagnostic use rather than for stereometric measurement, there is no direct means of obtaining a precise statement of the system geometry necessary for this application. Therefore, stereoradiographs of a special calibration device are prepared. This device consists of thirteen lead markers set at known positions in a radiolucent Plexiglas™ prism. The prism is placed so that at least eight of the markers will be visible in each of the two X-ray exposures. The body coordinates of this prism as it is radiographed become the arbitrary laboratory reference frame. Two X-rays are taken, an anterior-posterior and a lateral projection. Markers placed on the auditory meatus and the notches of the inferior orbital ridges, as well as those in the Plexiglas™ T-plate, are clearly visible in them. The positions of these images in each of the projections can then be made to yield their lab-oriented positions, which will in turn yield the orientations and lab-oriented positions of the anatomical and instrumentation coordinates, and finally the transformation from one to the other.

The anterior-posterior and lateral X-rays are also used to quantize the T-1 instrumentation coordinate data. Markers placed on the suprasternal notch and over the T-1 posterior spinous process, as well as those in the plexiglass T-plate, are clearly visible in each X-ray. In addition, the superior corners of the right and left articular facets are visible in the anterior-posterior X-rays, and the anterior-superior corner and the superior and inferior corners of the posterior spinous process are visible in the lateral ones.

After X-rays have been taken and developed, they are examined to determine that all the various markers and anatomical features are visible. The person overseeing radiological operations locates and marks the various anatomical landmarks for digitization. In digitization, the position of each marker and landmark is measured and compared to the orientation information. Digitized positions are subjected to computational procedures to determine, first, the geometrical layout of the radiologic equipment, and then the geometry relating the mounts to the anatomy. The computation is essentially a least squares iteration.

## *X-Ray Anthropometry Transformation Program*

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### 2. FUNCTION

The following program and subroutines are used:

- MXRAY: Main program, which allows operator to select major options.
- AXEB: Inverts a matrix.
- CDNTPT: Calculates and prints transformation matrices.
- POBKGS: Calculates the likeliest position and orientation of a rigid body of known geometry.
- PXRY2: Calculates the least-squared coordinates.
- RLSWCH: Right-left switch.
- OPTION: Compares calculations and uses switched values if indicated.
- PRISM2: Reads and reduces calibration data.
- HTRANS: Calculates the position and orientation of the head coordinates.

### 3. MAIN PROGRAM

#### 3.1 MXRAY

This interactive program calculates X-ray anthropometry transformation matrices. The output consists of a printout and a data file containing the "Instrumentation vs. Anatomy" matrix, which is the input to the Anthropometry Data Base Update program. The following input is requested:

Subject Number (A6 format)  
Date of X-ray (MON YR A6 format)  
Julian Date of X-ray (I5 format)  
Pcode (*a* for A-plate or *t* for T-plate, A1 format)  
Julian Date of X-ray (day, year; I3,I2 format)  
Option code "jo" and processing code "kk" (2I1 format)  
    jo = 1 exercise the option (standard  
        operation — always 1)  
    jo = (any other number) option not exercised  
    kk = 1 process head and neck data  
    kk = 2 process head data only  
    kk = 3 process neck data only  
    kk = 4 process pelvic data only  
    kk = 7 redo program  
    kk = 9 end program  
Comments for head data

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

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Comments for neck data  
Block or No block for neck data

### 3.2 COMPILATION

All of the subroutines needed to execute MXRAY are stored in the file 'libxrayantm.a,' which is in the users' library ('/usr/lib/'). The program was compiled with the following command:

```
fc mxray.f /usr/libxrayantm.a -o mxray
```

### 3.3 DATA FILES

The file 'input' should always contain the subject data to be processed. The file 'output' will always be used to store the processed data. The file 'update' will always be used to store the transformation matrices. The file 'IOupdate' is a shell procedure that updates the input file, saves the previous output, and executes the main program.

### 3.4 EXECUTION

The executable code is stored in the file 'mxray.' To execute the program, execute the shell procedure 'IOupdate' by typing the appropriate file names in the following form:

***IOupdate newinputfilename oldoutputfilename oldupdatefilename***

Assuming subject H00290 was processed, the output from program execution is in files 'output' and 'update.' The data will be saved on the next execution of 'IOupdate.' The user input would be as follows:

***IOupdate input H00290.out H00290.up***

The file 'input' contains data for the subject to be processed. The naming convention for the files 'oldoutputfilename' and 'oldupdatefilename' is 'subjectID.out' and 'subjectID.up,' respectively.

The program will request input, organize and coordinate calculations. To get printed results, execute the "lp" command using the file 'output.'

## X-Ray Anthropometry Transformation Program

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### 4. SUBROUTINE DESCRIPTIONS

#### 4.1 AXEB

The routine AXEB inverts a matrix. The call is:

*call axeb (a,n,n1,jc)*

where

A = the matrix to be inverted  
N = the number of rows in matrix A  
N1 = the number of columns in matrix A  
JC = the work array used by this routine

#### 4.2 CDNTPT

The routine CDNTPT calculates in anatomical coordinates the vector from the instrument origin to the anatomical origin. It also generates the transformation matrix that translates a vector from the instrumentation coordinate system to the anatomical coordinate system. The call is:

*call cdntpt*

#### 4.3 POBKKG

The routine POBKKG calculates the vector defining the instrument origin lab coordinates and also calculates the transformation matrix from the instrument to the laboratory system. The call is:

*call pobkg (x,xx,ii,pp,a)*

where

X = represents a two-dimensional, three-by-three array containing the best least squares coordinates of the center right, and left T-plate BBs in the lab coordinate system as calculated in subroutine PXRY2  
XX = a two-dimensional, three-by-three array containing the location of the three instrument BBs in the instrument coordinate system  
II = the number of BBs on the T-plate (normally 3)  
PP = the vector defining the instrument origin in lab coordinates

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

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A = the transformation matrix ( $3 \times 3$ ) that takes a vector from the instrument coordinate system and translates it into the lab coordinate system

### 4.4 PXRY2

The routine PXRY2 calculates the best least square coordinates in the lab coordinate system. The call is:

*call pxry2 (x,a,r,xr)*

where

X = the best least squares estimate of BBs in lab coordinates  
A = the measured X-ray coordinates of the nth BB  
R = the sum of the square of the error between the measured and the projected X-ray coordinates  
XR = the X-ray coordinates obtained by projection of the calculated position of the BB onto the X-ray film planes

### 4.5 RLSWCH

The routine RLSWCH allows an optional right-left switch of the anterior-posterior (AP) and lateral X-ray coordinates. The call is:

*call rlsrch (j,k,sp)*

where

J = represents the index of the right side BB on the lateral view as obtained from measurement  
K = the index of the right side BB so labeled by this subroutine and has identical X-ray coordinates as the measured labeled left side BB  
SP = the array containing the measured coordinates of the AP and lateral views. This array also contains the right-left/left right switch of coordinates in the last 24 elements

## X-Ray Anthropometry Transformation Program

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### 4.6 OPTION

The routine OPTION compares the error associated with the measured location of right and left BBs with the error associated with the right and left BBs as defined in subroutine RLSWCH. If the error is smaller for the point as labeled, the routine prints "Option Not Indicated." If the error is greater, the routine may either disregard the as-labelled calculation and use that of the switched labels (defined by subroutine RLSWCH) and print "Option Indicated and Exercised"; or, if the operator wishes, continue to use the as-labelled calculation and print "Option Indicated But Ignored." The call is:

*call option (j,k,er,x,jo)*

where

- J = the number of the right-side BBs as obtained from lateral view measurements
- K = the number of the right-side BBs as labeled by subroutine RLSWCH
- ER = a one-dimensional array containing the sum of the squares of the difference between the "measured" and "calculated" AP and lateral X-ray coordinates of the BB under scrutiny
- X = two-dimensional, three-by-thirteen array containing the best least squares coordinates for the BB under scrutiny, as calculated in subroutine PXRY2
- JO = the code that determines if an option is to be exercised, as follows:
  - 1 — Exercise option
  - Any other number — Do no exercise option

### 4.7 PRISM2

The routine PRISM2 reads and reduces calibration data. This subroutine establishes the lab coordinate system and calculates both the AP and lateral camera orientations and positions. The call is:

*call prism2 (xop,xsp,cp)*

where

- XOP = the best estimate of the vector from lab to X-ray origin in X-ray coordinates
- XSP = the best estimate of the vector from source origin to X-ray origin in X-ray coordinates

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

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CP = the best estimate of the transformation matrix  
from lab to X-ray

### 4.8 HTRANS

The routine HTRANS calculates the anatomical origin of the head in lab coordinates and the transformation matrix, which takes a vector in lab coordinates and transforms it into a vector in the head anatomical coordinate system. The call is:

*call htrans (x)*

where

X = the array containing the lab coordinates of the four head anatomical BBs plus the three T-plate BBs plus the six coordinates obtained by performing a right-left/left-right switch of measure-defined locations of BBs from the X-ray view

## 5. UTILITY SUBROUTINES

Two utility subroutines are used in the X-ray transformation program and are included for completeness. A brief description of each is given in this section.

### 5.1 CAPS

The routine CAPS enables the 'caps' mode on the HP-2627A terminal. The call is:

*call caps*

### 5.2 CAPOFF

The routine CAPOFF disables the 'caps' mode on the HP-2627A terminal. The call is:

*call capoff*

## *X-Ray Anthropometry Transformation Program*

---

### **REFERENCE**

Becker, E. B., "Stereoradiographic Measurements for Anatomically Mounted Instruments," *Proceedings of the Twenty-First STAPP Car Crash Conference*, Society of Automotive Engineers, Inc., Warrendale, PA, pp. 477-505, October 1977.

## **APPENDIX A**

### **Program Listings**

## X-Ray Anthropometry Transformation Program

---

FORTRAN 77/UX HP92430A.07.04 COPYRIGHT HEWLETT-PACKARD CO. 1983.  
Thu Jul 26 14:20:18 1990

```
1      subroutine axeb (a,n,n1,jc)
2  c
3  c This routine inverts a matrix
4  c
5  c a - matrix to be inverted
6  c n - number of rows in matrix a
7  c n1 - number of columns in matrix a
8  c jc - work array used by this routine
9  c
10     real a(n,n1)
11     integer jc(n)
12  c
13  c assign a unit number to the output file
14     open(11,file='output')
15  c
16     do 70 i=1,n
17       x=-1.
18       m=i
19       do 10 j=m,n
20         if (abs(a(j,i)).lt.x) go to 10
21         x=abs(a(j,i))
22         l=j
23   10 continue
24       c
25         c check matrix for singularity
26         c
27         jc(i)=114
28         if (x) 20,90,20
29   20 do 30 j=1,n1
30     x=a(i,j)
31     a(i,j)=a(l,j)
32   30 a(l,j)=x
33     x=a(i,i)
34     a(i,i)=1.
35     do 40 j=1,n1
36   40 a(i,j)=a(i,j)/x
37     do 60 j=1,n
38       if (j.eq.i) go to 60
39       x=a(j,i)
40       a(j,i)=0.
41       do 50 k=1,n1
42   50 a(j,k)=a(j,k)-x*a(i,k)
43   60 continue
44   70 continue
45     do 80 i3=1,n
46       i=1+n-i3
47       l=jc(i)
48       do 80 j=1,n
49         x=a(j,i)
50         a(j,i)=a(j,l)
51   80 a(j,l)=x
52     return
53   90 write (11,100)
54     jc(1)=-1
55     return
56  c
57    100 format (' disaster - ill conditioned matrix')
```

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

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58 c  
59 end

NUMBER OF ERRORS = 0 NUMBER OF WARNINGS = 0

## *X-Ray Anthropometry Transformation Program*

---

FORTRAN 77/UX HP92430A.07.04 COPYRIGHT HEWLETT-PACKARD CO. 1983.  
Thu Jul 26 14:20:19 1990

```
1      subroutine capoff
2  c
3  c  FUNCTION:
4  c  Disables the 'caps' mode on the HP-2627A terminal.
5  c
6  c  BY:
7  c  D. Francis
8  c  Naval Biodynamics Laboratory
9  c  New Orleans, Louisiana
10 c  28 June 1988
11 c
12     equivalence (ESCA,IEA),(AK0,IK0),(PP,IPP)
13 c
14     character*2 ESCA,AK0,PP
15 c
16     data IESCA/o'015446'/,K0/o'065460'/,IPSP/o'050040'/
17 c
18     call mvbits(IESCA,0,16,IEA,16)
19     call mvbits(K0,0,16,IK0,16)
20     call mvbits(IPSP,0,16,IPP,16)
21 c
22     write(6,10) ESCA,AK0,PP
23 10  format(6a2)
24 c
25     return
26     end
```

NUMBER OF ERRORS = 0 NUMBER OF WARNINGS = 0

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

---

FORTRAN 77/UX HP92430A.07.04 COPYRIGHT HEWLETT-PACKARD CO. 1983.  
Thu Jul 26 14:20:19 1990

```
1      subroutine caps
2  c
3  c  FUNCTION:
4  c  Enables the 'caps' mode on the HP-2627A terminal.
5  c
6  c  BY:
7  c  D. Francis
8  c  Naval Biodynamics Laboratory
9  c  New Orleans, Louisiana
10 c  28 June 1988
11 c
12     equivalence (ESCA,IEA),(AK1,IK1),(PP,IPP)
13 c
14     character*2 ESCA,AK1,PP
15 c
16     data IESCA/o'015446'/,K1/o'065461'/,IPSP/o'050040'/
17 c
18     call mvbits(IESCA,0,16,IEA,16)
19     call mvbits(K1,0,16,IK1,16)
20     call mvbits(IPSP,0,16,IPP,16)
21 c
22     write(6,10) ESCA,AK1,PP
23 10  format(6a2)
24 c
25     end
```

NUMBER OF ERRORS = 0 NUMBER OF WARNINGS = 0

## X-Ray Anthropometry Transformation Program

---

FORTRAN 77/UX HP92430A.07.04 COPYRIGHT HEWLETT-PACKARD CO. 1983.  
Thu Jul 26 14:20:24 1990

```
1      subroutine cdntpt
2  c
3  c This subroutine calculates in anatomical coordinates the vector from
4  c instrument origin to the anatomical origin. It also generates the
5  c transformation matrix that take a vector from the instrumentation
6  c coordinate system to the anatomical coordinate system.
7  c
8  c input consist of common/blk1/arrays;
9  c
10 c   xh(3) = created in subroutine htrans and is the vector defining the
11 c   anatomical origin in lab coordinates.
12 c
13 c   ah(3,3) = created in the subroutine htrans and is the transformation
14 c   matrix that takes a vector from the lab into the anatomical
15 c   coordinate system.
16 c
17 c
18 c   xt(3) = created in subroutine pobkg where it was called 'pp(i)';
19 c   vector defining the instrument origin in lab coordinates
20 c
21 c   at(3,3) = created in subroutine pobkg where it was called 'a(3,3)';
22 c   transformation matrix that takes a vector from instruments
23 c   into the lab coordinate system.
24 c
25 c Output consists of arrays
26 c
27 c   r(i) = r(i) + ah(i,j) * xh(k,j)
28 c   The instrument origin vector transformed into the head
29 c   anatomical coordinate system where
30 c   xh(k,j) = xt(i)-xh(i) the vector from the
31 c   instrument origin to the anatomy origin in lab coordinates.
32 c
33 c   aa(i,j) = aa(i,j) + ah(i,k)*at(k,j)
34 c   where aa(i,j) = the transformation matrix that takes a
35 c   vector in instrument coordinates and expresses it in
36 c   anatomy coordinates.
37 c
38 c
39      real aa(3,3), r(3)
40      integer sday,syear,eday,eyear
41      character*6 nsub,dat,blk
42      character*6 not(9)
43 c
44      common /blk1/ xh(3),ah(3,3),xt(3),at(3,3)
45      common /c1/ nsub,nmnt,dat,blk,not,sday,syear,eday,eyear
46 c
47 c assign a unit number to the output file
48      open(11,file='output')
49      open(12,file='update')
50 c
51      write (11,100)
52      do 10 i=1,3
53      write (11,50) xh(i),(ah(i,j),j=1,3),xt(i),(at(i,j),j=1,3)
54 10  xh(i)=xt(i)-xh(i)
55      write (11,90)
56      do 30 i=1,3
57      r(i)=0.
58      do 20 j=1,3
```

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

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```
59      aa(i,j)=0.
60      r(i)=r(i)+ah(i,j)*xh(j)
61      do 20 k=1,3
62 20  aa(i,j)=aa(i,j)+ah(i,k)*at(k,j)
63 30  write (11,60) r(i),(aa(i,j),j=1,3)
64      c
65      c   write output data to update file
66      c
67      write(12,110) nsub,nmnt,dat,blk,(not(j),j=1,5),sday,s year,eday,
68 *eyear
69      write(12,80) (r(i),i=1,3)
70      do 40 i=1,3
71      write(12,70) (aa(j,i),j=1,3)
72 40  continue
73      c
74      return
75 c
76 50 format (2(5x,f10.4,3x,3f10.8))
77 60 format (5x,f10.4,3x,3f10.8)
78 70 format (3(f10.8,2x))
79 80 format (3(f10.6,2x))
80 90 format (///12x,'instrumentation vs anatomy')
81 100 format (///18x,'anatomy vs lab',27x,'instrumentation vs lab')
82 110 format (a6,2x,i4,2x,2(a6,3x),5a6,4x,2(i3,i4))
83 c
84      end
```

NUMBER OF ERRORS = 0 NUMBER OF WARNINGS = 0

## X-Ray Anthropometry Transformation Program

---

FORTRAN 77/UX HP92430A.07.04 COPYRIGHT HEWLETT-PACKARD CO. 1983.  
Thu Jul 26 14:20:25 1990

```
1      subroutine htrans (x)
2  c
3  c subroutine htrans
4  c
5  c
6  c This subroutine calculates the anatomical origin of the head
7  c in lab coordinates and the transformation matrix which takes
8  c a vector in lab coordinates and transforms it into a vector
9  c in the head anatomical coordinate system.
10 c
11 c input consist of:
12 c x(3,13) = the lab coordinates of the 4 head anatomical bee bee's
13 c plus t3 t-plate bee bee's plus the 6 coordinates obtained
14 c by performing a right-left/left-right switch of measure
15 c defined locations of bee's bee's from the x-ray view.
16 c These are computed in subroutine pxryz by a least squares
17 c technique.
18 c
19 c xt(3t(3) = created in subroutine pobkg where it was called 'pp(i)'
20 c and and is the vector defining the instrument origin i
21 c lab coordinates.
22 c
23 c at(3,3) = created in subroutine 'pobkg' where it was called 'a(3,3)'
24 c and is the transformation matrix that takes a vector from
25 c instrument trument into the lab coordinates system.
26 c
27 c output consist of:
28 c xh(i) = (x(i,2) + x(i,1))/2 = anat origin in lab.
29 c
30 c ah(3,3) = is the transformation matrix which takes a vector in lab
31 c coordinates and transforms it into a vector in the head
32 c anatomical system.
33 c
34     common /blk1/ xh(3),ah(3,3),xt(3),at(3,3)
35     real x(3,13)
36     do 10 i=1,3
37     xh(i)=(x(i,2)+x(i,1))/2.
38     ah(1,i)=x(i,3)+x(i,4)-x(i,1)-x(i,2)
39     ah(2,i)=x(i,2)+x(i,4)-x(i,1)-x(i,3)
40     10 ah(3,i)=0.
41     do 20 i=1,3
42     ah(3,1)=ah(3,1)+ah(1,i)**2
43     ah(3,2)=ah(3,2)+ah(2,i)**2
44   20 ah(3,3)=ah(3,3)+ah(1,i)*ah(2,i)
45     ah(3,1)=sqrt(ah(3,1))
46     ah(3,2)=sqrt(ah(3,2))
47     ah(3,3)=ah(3,3)/(ah(3,1)*ah(3,2))
48     r=1./(sqrt(1.-ah(3,3)**2)*ah(3,2))
49     do 30 i=1,3
50     ah(1,i)=ah(1,i)/ah(3,1)
51   30 ah(2,i)=r*(ah(2,i)-ah(3,2)*ah(3,3)*ah(1,i))
52     ah(3,1)=ah(1,2)*ah(2,3)-ah(1,3)*ah(2,2)
53     ah(3,2)=ah(2,1)*ah(1,3)-ah(1,1)*ah(2,3)
54     ah(3,3)=ah(1,1)*ah(2,2)-ah(1,2)*ah(2,1)
55     return
56 c
57 end
```

NUMBER OF ERRORS = 0 NUMBER OF WARNINGS = 0

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

FORTRAN 77/UX HP92430A.07.04 COPYRIGHT HEWLETT-PACKARD CO. 1983.  
Thu Jul 26 14:20:26 1990

```
1  c  main and zz03 updated 9/4/80 by p. shimp to handle
2  c  processing of a-plates
3  c
4      common /blk1/ xh(3),ah(3,3),xt(3),at(3,3)
5      common /blk2/ xo(3,2),xs(3,2),c(3,3,2)
6      common /c1/ nsub,nmmt,dat,blk,not,sday,syear,eday,eyear
7  c
8      character*6 ksubj,nsub,dat,blk
9      character*6 not(9)
10     character*1 pcode
11     real w(3), p(4), pp(3,4)
12     integer jp(3)
13     real xp(2,2), x(3,13), xx(3)
14     real xtp(4)
15     real sp(2,2,13), er(13)
16     real xk(3,3), x1(3,3), x2(3,3)
17     real xpn(6)
18  c
19  c  the variables day and year refer to date of x-ray
20  integer day,year
21  c  The variables sday, syear, eday & eyear
22  c  are the start date and end date for the data.
23  integer date,sday,syear,eday,eyear
24  c
25      real spi(4,13)
26      equivalence (sp(1,1,1),spi(1,1))
27  c
28      data x1 /0.,0.,0.,6.286,-6.286,0.,6.286,6.286,0./
29      data x2 /-.3556,0.,.175,5.9304,-6.286,.175,5.9304,6.2 86,.175/
30  c
31  c  assign a unit number to the input and output files
32      open(10,file='input')
33      open(11,file='output')
34  c
35 10 continue
36  c
37  c  put terminal in 'caps' mode.....
38      call caps
39  c
40  c  request user input.....
41  c
42      write(6,*) 'enter subject number (a6 format)'
43      read(5,536) nsub
44      write(6,*) 'enter date of xray (MON YR a6 format)'
45      read(5,536) dat
46      write(6,*) 'enter Julian date of xray (i5 format)'
47      read(5,565) idate
48  c
49  c.....read data from input file .....
50  c
51      read(10,535) ksubj,date,mmount,nmount,sday,syear,eday ,eyear
52  c
53  c
54  c.....check for correct input data.....
55      if(ksubj.ne.nsub) go to 525
56      if(date.ne.idate) go to 12
57      go to 18
58 12 write(6,15)
```

## X-Ray Anthropometry Transformation Program

---

```
59 15 format('Dates do not match. Enter Y to continue, N to stop.')
60      read(5,540) pcode
61      if(pcode.ne.'Y') go to 530
62 18 continue
63 c
64 c.....initialize the variable 'blk'
65     blk=' '
66 c
67     c.....process the data.....
68 c
69     write(6,*) 'enter pcode (A for a-plate or T for t-plate)'
70     read (5,540) pcode
71     if (PCODE.NE.'A'.AND.PCODE.NE.'T') PRINT 550
72     if (PCODE.NE.'A'.AND.PCODE.NE.'T') STOP 1
73     write(6,*) 'enter xray Julian date'
74     read (5,545) day,year
75     d=year*1000+day
76     do 30 i=1,3
77     do 30 j=1,3
78     if (PCODE.EQ.'A') go to 20
79     XK(i,j)=X1(i,j)
80     go to 30
81 20 XK(i,j)=X2(i,j)
82 30 continue
83     if (PCODE.NE.'A') go to 35
84 c in the event x-ray was taken after jan 1,1981 change z component
85 c to.238 for A plates only
86     if (d.lt. 81001) go to 35
87     do 32 i=1,3
88     XK(3,i)=.238
89     32 continue
90 35 call prism2 (xo,xs,c)
91 40 write(6,555)
92     read(5,560) jo,kk
93     if(kk.eq.9) go to 530
94     if(kk.eq.7) go to 10
95     if (kk.eq.3) go to 220
96     if (kk.eq.4) go to 450
97 c
98 c.....mouth data calculations
99 c
100    write(6,*) 'enter comments for head data (9a6 format)'
101    read(5,575) not
102    nmnt=mmount
103    write (11,580)
104    write (11,590) nsub,nmnt,dat,(not(j),j=1,9)
105    write(11,630)
106    write(11,640)
107    write(11,650)
108    write(11,660)
109    write(11,820)
110    write(11,670)
111 c
112    do 50 j=1,4
113    read(10,620) (spi(j,i),i=1,7)
114    do 50 i=1,7
115    spi(j,i)=spi(j,i)*2.54
116 50 continue
117 c
118 c interchange ltp and rtp coordinates when processing a-plate
119 if (PCODE.NE.'A') go to 70
120 do 60 j=1,4
```

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

```
121      t=spi(j,6)
122      spi(j,6)=spi(j,7)
123      spi(j,7)=t
124 60  continue
125 70  continue
126      call rlsrch (1,8,sp)
127      call rlsrch (3,10,sp)
128      call rlsrch (6,12,sp)
129      do 210 i=1,13
130      call pxry2 (x(1,i),sp(1,1,i),er(i),xtp)
131      if (i.eq.1) go to 80
132      if (i.eq.2) go to 90
133      if (i.eq.3) go to 100
134      if (i.eq.4) go to 110
135      if (i.eq.5) go to 120
136      if (i.eq.6) go to 130
137      if (i.eq.7) go to 140
138      if (i.eq.8) go to 150
139      if (i.eq.9) go to 160
140      if (i.eq.10) go to 170
141      if (i.eq.11) go to 180
142      if (i.eq.12) go to 190
143      if (i.eq.13) go to 200
144 80  write(11,680) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
145      go to 210
146 90  write(11,690) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er()
147      go to 210
148 100 write(11,700) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
149      go to 210
150 110 write(11,710) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
151      go to 210
152 120 write(11,720) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
153      go to 210
154 130 write(11,730) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
155      go to 210
156 140 write(11,740) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
157      go to 210
158 150 write(11,750)
159      write(11,760) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
160      go to 210
161 160 write(11,770) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
162      go to 210
163 170 write(11,780) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
164      go to 210
165 180 write(11,790) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
166      go to 210
167 190 write(11,800) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
168      go to 210
169 200 write(11,810) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
170 210 continue
171      call option (1,8,er,x,jo)
172      call option (3,10,er,x,jo)
173      call option (6,12,er,x,jo)
174      if (PCODE.eq.'A') write(11,850)
175      call pobkg (x(1,5),xk,3,xt,at)
176      call htrans (x)
177      write (11,580)
178      write (11,600) nsub,nmnt,dat
179      write (11,610) (not(j),j=1,9)
180      call cdntpt
181      if (kk.eq.2) go to 40
182 c
```

## X-Ray Anthropometry Transformation Program

---

```
183 c.....neck data calculations
184 c
185 220 continue
186     nmmt=nmount
187     write(6,*) 'enter comments for neck data (9a6 format)'
188     read(5,575) not
189     write(6,*) 'enter block or noblock (a6 format)'
190     read(5,536) blk
191     write (11,860)
192     write (11,600) nsub,nmmt,dat
193     write (11,610) (not(j),j=1,9)
194 c
195 c.....read the required data
196 c
197     do 230 j=1,4
198     read(10,620) (spi(j,i),i=1,5)
199     do 230 i=1,5
200     spi(j,i)=spi(j,i)*2.54
201 230 continue
202     call rlsrch (4,6,sp)
203     sc1=0.
204     sc2=0.
205     write(11,630)
206     write(11,640)
207     write(11,650)
208     write(11,660)
209     write(11,820)
210     write(11,670)
211     do 310 i=1,7
212     call pxry2 (x(1,i),sp(1,1,i),er(i),xtp)
213     if (i.gt.5) sc2=sc2+sp(2,2,i)-xtp(4)
214     if (i.eq.4.or.i.eq.5) sc2=sc2-sp(2,2,i)+xtp(4)
215     if (i.lt.6) sc1=sc1+sp(2,2,i)-xtp(4)
216     if (i.eq.1) go to 240
217     if (i.eq.2) go to 250
218     if (i.eq.3) go to 260
219     if (i.eq.4) go to 270
220     if (i.eq.5) go to 280
221     if (i.eq.6) go to 290
222     if (i.eq.7) go to 300
223 240 write(11,870) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
224     go to 310
225 250 write(11,880) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
226     go to 310
227 260 write(11,890) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
228     go to 310
229 270 write(11,900) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
230     go to 310
231 280 write(11,910) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
232     go to 310
233 290 write(11,750)
234     write(11,920) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
235     go to 310
236 300 write(11,930) (spi(j,i),j=1,4),(x(ii,i),ii=1,3),er(i)
237 310 continue
238     call option (4,6,er,x,jo)
239     ai=er(4)**2+er(5)**2-er(6)**2-er(7)**2
240     if (ai.gt.0.) sc1=sc1+sc2
241     call pobkg (x(1,3),xk,3,xt,at)
242     read(10,620) xp
243     do 320 i=1,2
244     do 320 j=1,2
```

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

---

```
245      xp(i,j)=xp(i,j)*2.54
246      320  continue
247      write (11,940) xp
248      ah(1,1)=-xs(3,1)*(xp(2,2)-xp(2,1))
249      ah(1,2)=xs(3,1)*(xp(1,2)-xp(1,1))
250      ah(1,3)=(xs(1,1)-xp(1,1))*(xp(2,2)-xp(2,1))-(xs(2,1)-xp(2,1))*(xp(
251      11,2)-xp(1,1))
252      do 330 i=1,3
253      xx(i)=x(i,2)-x(i,1)
254      ah(3,i)=0.
255      do 330 j=1,3
256      330 ah(3,i)=c(j,i,1)*ah(1,j)+ah(3,i)
257      ah(2,1)=ah(3,2)*xx(3)-ah(3,3)*xx(2)
258      ah(2,2)=ah(3,3)*xx(1)-ah(3,1)*xx(3)
259      ah(2,3)=ah(3,1)*xx(2)-ah(3,2)*xx(1)
260      ai=0.
261      do 340 i=1,3
262 340  ai=ai+ah(2,i)**2
263      ai=sqrt(ai)
264      do 350 i=1,3
265      1 350 ah(2,i)=ah(2,i)/ai
266 c
267      read(10,620) (xpn(k),k=1,6)
268      xp(1,1)=xpn(1)
269      xp(2,1)=xpn(2)
270      xp(1,2)=(xpn(3)+xpn(5))/2.0
271      xp(2,2)=(xpn(4)+xpn(6))/2.0
272 c
273      do 360 i=1,2
274  do 360 j=1,2
275      xp(i,j)=xp(i,j)*2.54
276 360  continue
277      write (11,960) xp
278      do 370 ill=1,2
279      370 xp(2,ill)=xp(2,ill)-sc1/5.
280      g1=0.
281      do 390 i=1,3
282      g3=x(i,1)
283      xx(i)=0.
284      do 380 j=1,3
285      xx(i)=xx(i)+ah(2,j)*c(i,j,2)
286 380  g3=g3+c(j,i,2)*(xo(j,2)-xs(j,2))
287 390  g1=g1+g3*ah(2,i)
288      do 420 i=1,2
289      ga=g1/((xp(1,i)-xs(1,2))*xx(1)+(xp(2,i)-xs(2,2))*xx(2)-xs(3,2)*x
290      1x(3))
291      do 400 j=1,2
292 400  x(j,8)=ga*(xp(j,i)-xs(j,2))+xs(j,2)
293      x(3,8)=(1.-ga)*xs(3,2)
294      l=i+5
295      do 410 j=1,3
296      x(j,l)=0.
297      do 410 k=1,3
298 410  x(j,1)=(x(k,8)-xo(k,2))*c(k,j,2)+x(j,1)
299 420  write (11,970) (x(j,1),j=1,3)
300      ai=0.
301      do 430 i=1,3
302      xh(i)=x(i,6)
303      ah(1,i)=xh(i)-x(i,7)
304 430  ai=ai+ah(1,i)**2
305      ai=sqrt(ai)
306      do 440 i=1,3
```

## X-Ray Anthropometry Transformation Program

---

```
307      440 ah(1,i)=ah(1,i)/ai
308      ah(3,1)=ah(1,2)*ah(2,3)-ah(1,3)*ah(2,2)
309      ah(3,2)=ah(2,1)*ah(1,3)-ah(1,1)*ah(2,3)
310      ah(3,3)=ah(1,1)*ah(2,2)-ah(1,2)*ah(2,1)
311      write (11,860)
312      write (11,600) nsub,nmnt,dat
313      write (11,610) (not(j),j=1,9)
314      call cdntpt
315      go to 40
316 c
317 c.....pelvic data calculations
318 c
319 450 continue
320      write (11,980)
321
322      do 460 k=1,4
323 460 pp(j,k)=0.
324 470 do 480 j=1,4
325 480 read (10,620) (spi(j,i),i=1,6)
326      do 490 i=1,4
327      write (11,840) i,(spi(j,i),j=1,4)
328      call pxry2 (x(1,i),sp(1,1,i),er(i),xtp)
329 490 write (11,830) (x(ii,i),ii=1,3),er(i)
330      call htrans (x)
331      do 520 i=1,2
332      p(1)=(sp(2,i,6)-sp(2,i,5))*xs(3,i)
333      p(2)=(sp(1,i,5)-sp(1,i,6))*xs(3,i)
334      p(3)=(sp(1,i,5)-xs(1,i))*(sp(2,i,6)-xs(2,i))-(sp(2,i,5)-xs(2,i))
335      1*(sp(1,i,6)-xs(1,i))
336      p(4)=0.
337      do 500 j=1,3
338      p(4)=p(4)+p(j)*(xs(j,i)-xo(j,i))
339      w(j)=0.
340      do 500 k=1,3
341 500 w(j)=w(j)+p(k)*c(k,j,i)
342      do 510 j=1,3
343      p(4)=p(4)-w(j)*xh(j)
344      p(j)=0.
345      do 510 k=1,3
346 510 p(j)=p(j)+ah(j,k)*w(k)
347      write (11,990) p
348      do 520 k=1,3
349      do 520 j=1,4
350 520 pp(k,j)=pp(k,j)+p(k)*p(j)
351      read (5,1000) jj
352      if (jj.ne.0) go to 470
353      call axebs(pp,3,4,jp)
354      write (11,1010) (pp(j,4),j=1,3)
355      go to 40
356 c
357 c.....Output error message.....
358 c
359 525 write(6,*) 'Incorrect Subject Id - Processing Stopped'
360 c
361 530 call capoff
362      stop
363 c
364 535 format(a6,1x,i5,1x,2(i4,1x),2(i3,i4))
365 536 format(a6)
366 540 format (a1)
367 545 format(i3,i2)
368 550 format (1x,'plate code invalid, check first input card'/1x,'it
```

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

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```

369      1should contain an a or t in column 1')
370 555 format('enter option code jo and processing code kk (2i1) format'
371      **'jo = 1 to exercise the option (standard operation -always 1)''/
372      **'jo = any other number not to exercise the option'/
373      **'kk = 1 process head and neck data'/
374      **'kk = 2 process head data only'/
375      **'kk = 3 process neck data only'/
376      **'kk = 4 process pelvic data only'/
377      **'kk = 7 redo program'/
378      **'kk = 9 end program')
379 560 format (2i1)
380 565 format (i5)
381 570 format (a6,i4,10a6)
382 575 format (9a6)
383 580 format ('1 head anatomical to tee-plate transformation ')
384 590 format (' subject ',a6,' mount ',i4,' date ',a6,' notes '
385     19a6)
386 600 format (' subject ',a6,' mount ',i4,' date ',a6)
387 610 format (10x,' notes ',9a6)
388 620 format (7(f7.3,1x))
389 630 format (1h0,'bee bee id i measured x-ray coordinates
390     1 i lab coordinates i sum of the squares of the')
391 640 format (12x,'i',42x,'i',31x,'i difference between the')
392 650 format (12x,'i',42x,'i',31x,'i measured and best estimate')
393 660 format (12x,'i',8x,'ap',19x,'lat',10x,'i',31x,'i of x-ray coordin
394     ates')
395 670 format (1h0,11x,'i',4x,'x',10x,'y',10x,'x',8x,'y',6x'i',4x,'x',9x
396     1,'y',9x,'z',6x,'i',4x,'e')
397 680 format (1h,1x,'1 r.audit ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,
398     12x,f8.4,2x,f8.4,4x,f8.4)
399 690 format (2x,'2 l.audit ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,f
400     18.4,2x,f8.4,4x,f8.4)
401 700 format (2x,'3 r.orbit ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,f
402     18.4,2x,f8.4,4x,f8.4)
403 710 format (2x,'4 l.orbit ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,f
404     18.4,2x,f8.4,4x,f8.4)
405 720 format (2x,'5 c.t-plt ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,f
406     18.4,2x,f8.4,4x,f8.4)
407 730 format (2x,'6 r.t-plt ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,f
408     18.4,2x,f8.4,4x,f8.4)
409 740 format (2x,'7 l.t-plt ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,f
410     18.4,2x,f8.4,4x,f8.4)
411 750 format (1h0,14x,'right/left switch follow below')
412 760 format (1h0,1x,'8 r.audit ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,
413     12x,f8.4,2x,f8.4,4x,f8.4)
414 770 format (2x,'9 l.audit ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,f
415     18.4,2x,f8.4,4x,f8.4)
416 780 format (1x,'10 r.orbit ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,
417     1f8.4,2x,f8.4,4x,f8.4)
418 790 format (1x,'11 l.orbit ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,
419     1f8.4,2x,f8.4,4x,f8.4)
420 800 format (1x,'12 r.t-plt ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,
421     1f8.4,2x,f8.4,4x,f8.4)
422 810 format (1x,'13 l.t-plt ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,
423     1f8.4,2x,f8.4,4x,f8.4)
424 820 format (12x,'i',42x,'i',31x,'i')
425 830 format (5x,4f15.4)
426 840 format (1h0,2x,i2,4f15.4)
427 850 format (///1x,'note:when processing data from a-plate, ltp and rtp
428     1'1x,'are interchanged from the way in which they were entered to'
429     25x,'account for position of a-plate ')
430 860 format ('1 t-1 anatomical to tee-plate transformation ')

```

## X-Ray Anthropometry Transformation Program

---

```
431 870 format (1h0,1x,'1 p-spine ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,
432 12x,f8.4,2x,f8.4,4x,f8.4)
433 880 format (2x,'2 s-notch ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,f
434 18.4,2x,f8.4,4x,f8.4)
435 890 format (2x,'3 c.t-plt ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,f
436 18.4,2x,f8.4,4x,f8.4)
437 900 format (2x,'4 r.t-plt ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,f
438 18.4,2x,f8.4,4x,f8.4)
439 910 format (2x,'5 l.t-plt ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,f
440 18.4,2x,f8.4,4x,f8.4)
441 920 format (1h0,1x,'6 r.t-plt ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,
442 12x,f8.4,2x,f8.4,4x,f8.4)
443 930 format (2x,'7 l.t-plt ',f8.4,2x,f8.4,4x,f8.4,2x,f8.4,4x,f8.4,2x,f
444 18.4,2x,f8.4,4x,f8.4)
445 940 format (22h0 articular facets,4f10.4)
446 960 format (23h0 lateral projection,4f10.4)
447 970 format (5x,3f15.4)
448 980 format ('1 c.g. routine')
449 990 format (5x,4f14.6)
450 1000 format (i2)
451 1010 format (5x,3f14.6)
452 1020 format (a6,i5,a5,i2,2i7,20f7.3)
453 c
454     end
```

NUMBER OF ERRORS = 0 NUMBER OF WARNINGS = 0

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

FORTRAN 77/UX HP92430A.07.04 COPYRIGHT HEWLETT-PACKARD CO. 1983.  
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```
1      subroutine option (j,k,er,x,jo)
2  c
3  c      subroutine option
4  c
5  c
6  c      This subroutine compares the residual error associated with the
7  c      measured definition of the location of right and left bee bee's, with
8  c      the errors associated with subroutine rlsrch's definition of right
9  c      and left bee bee's. Finally it states the option with least error and
10 c      if told to do so will exercise the option.
11 c
12 c      The options are
13 c      1. option j vs. k not indicated
14 c      2. option j vs. k indicated but ignored
15 c      3. option j vs. k indicated and exercised
16 c
17 c      Input consist of
18 c      j = the number of the right-side bee bee's as obtained from
19 c            lateral view measurements
20 c
21 c      k = the number of the right-side bee bee's so labeled by
22 c            subroutine rlsrch has identical x-ray coordinates as the
23 c            measured labeled left side bee bee
24 c
25 c      er = the sum of the squares of the difference between the
26 c            'measured' and 'calculated' ap and lateral x-ray
27 c            coordinates of the bee bee under scrutiny.
28 c
29 c      x = calculated in subroutine pxryz and is the best least squares
30 c            lab coordinates for the bee bee under scrutiny.
31 c
32 c      jo = code which determines if an option is to be exercised.
33 c
34 c
35      real x(3,13),er(13)
36 c
37 c      assign a unit number to the output file
38      open(11,file='output')
39 c
40      r=er(k)*er(k)+er(k+1)*er(k+1)-er(j)*er(j)-er(j+1)*er(j+1)
41      if (r.gt.0.) go to 30
42      if (jo.ne.1) go to 20
43      write (11,40) j,k
44      l=j+1
45      do 10 i=1,3
46      x(i,j)=x(i,k)
47 10      x(i,l)=x(i,k+1)
48      return
49 20      write (11,50) j,k
50      return
51 30      write (11,60) j,k
52      return
53 c
54 40      format (9h option,i2,4h vs,i2,24h indicated and exercised)
55 50      format (9h option,i2,4h vs,i2,23h indicated but ignored)
56 60      format (9h option,i2,4h vs,i2,15h not indicated)
57 c
58      end
NUMBER OF ERRORS = 0 NUMBER OF WARNINGS = 0
```

## X-Ray Anthropometry Transformation Program

---

FORTRAN 77/UX HP92430A.07.04 COPYRIGHT HEWLETT-PACKARD CO. 1983.  
Thu Jul 26 14:20:57 1990

```
1      subroutine pobkg (x,xx,ii,pp,a)
2  c
3  c subroutine pobkg
4  c
5  c
6  c This subroutine calculates the vector defining the
7  c instrument origin pp lab coordinates and also calculates
8  c the transformation matrix [instr/lab]
9  c a(i,j) = at(i,j) of main program.
10 c
11 c input consist of:
12 c x(3,3) = best least squares coordinates of the center, right,
13 c           and left t-plate bee bee's in the lab coordinate
14 c           system as calculated in subroutine pxxyz
15 c
16 c ii      = 3 the number of bee bee's on the t-plate
17 c
18 c xx(3,3) = xk in main program and is the location of the 3
19 c           instrument bee bee's in the instrument coordinate system.
20 c
21 c output consist of:
22 c pp(i)   = vector defining the instrument origin in lab coordinates
23 c           also xt(i) in the main program and in subroutine cdnpt
24 c
25 c a(i,j)  = transformation matrix which takes a vector from the
26 c           instrument coordinate system and puts it in the lab
27 c           coordinate system. Also called at(j,j) in the main program
28 c
29     real xp(3)
30     real x(3,3), xx(3,3), pp(3), a(3,3), xm(3,3)
31 c
32 c assign a unit number to the output file
33 c open(11,file='output')
34 c
35     write(11,440)
36     write(11,120)
37     write(11,130)
38     write(11,140)
39     write(11,150)
40     write(11,160)
41     write(11,170)
42     write(11,180)
43     write(11,190)
44     write(11,200)
45     write(11,210)
46     write(11,220)
47     write(11,230)
48     write(11,240)
49     write(11,250)
50     write(11,260)
51     write(11,270)
52     write(11,280)
53     write(11,290)
54     write(11,300)
55     write(11,310)
56     write(11,320)
57     write(11,330)
58     write(11,340)
```

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

---

```
59      write(11,350)
60      write(11,360)
61      write(11,370)
62      write(11,380)
63      write(11,390)
64      write(11,400)
65      write(11,410)
66      write(11,420)
67      write(11,430)
68      write(11,110)  xx
69  c
70  c
71      do 20 i=1,3
72      xp(i)=0.
73      pp(i)=0.
74      do 10 j=1,ii
75      pp(i)=pp(i)+x(i,j)
76  10  xp(i)=xp(i)+xx(i,j)
77      xp(i)=xp(i)/float(ii)
78  20  pp(i)=pp(i)/float(ii)
79      do 40 i=1,3
80      do 30 j=1,3
81      a(i,j)=0.
82      xm(i,j)=0.
83      do 30 k=1,ii
84  30  xm(i,j)=xm(i,j)+(x(i,k)-pp(i))*(xx(j,k)-xp(j))
85  40  a(i,i)=1.
86      do 80 ijk=1,10
87      w=atan2((xm(3,2)-xm(2,3)),(xm(2,2)+xm(3,3)))
88      c=cos(w)
89      s=sin(w)
90      do 50 k=1,3
91      r=a(2,k)
92      a(2,k)=c*r-s*a(3,k)
93      a(3,k)=s*r+c*a(3,k)
94      r=xm(k,2)
95      xm(k,2)=c*r-s*xm(k,3)
96  50  xm(k,3)=s*r+c*xm(k,3)
97      w=atan2((xm(1,3)-xm(3,1)),(xm(1,1)+xm(3,3)))
98      c=cos(w)
99      s=sin(w)
100     do 60 k=1,3
101     r=a(1,k)
102     a(1,k)=c*r+s*a(3,k)
103     a(3,k)=c*a(3,k)-s*r
104     r=xm(k,1)
105     xm(k,1)=c*r+s*xm(k,3)
106  60  xm(k,3)=c*xm(k,3)-s*r
107     w=atan2((xm(2,1)-xm(1,2)),(xm(1,1)+xm(2,2)))
108     c=cos(w)
109     s=sin(w)
110     do 70 k=1,3
111     r=a(1,k)
112     a(1,k)=c*r-s*a(2,k)
113     a(2,k)=c*a(2,k)+s*r
114     r=xm(k,1)
115     xm(k,1)=c*r-s*xm(k,2)
116  70  xm(k,2)=c*xm(k,2)+s*r
117     r=(abs(xm(2,3)-xm(3,2))+abs(xm(1,2)-xm(2,1))+abs(xm(1,3)-xm(3,1)
118     1))/((xm(1,1)+xm(2,2)+xm(3,3)))
119     if (r.ge.0..and.r.lt..000001) go to 90
120  80  continue
```

## X-Ray Anthropometry Transformation Program

---

```
121 90 do 100 i=1,3
122      do 100 j=1,3
123 100 pp(i)=pp(i)-a(i,j)*xp(j)
124      return
125 c
126 110 format (///60x,'x y z'//48x,'bb 1 ',3f9.4//6x,'loca
127      ion of bee bees in instrumentation: bb 2 ',3f9.4//48x,'bb 3  ',
128      3f9.4)
129 120 format (//50x,'-----\'')
130 130 format (47x,'///',17x,'\'')
131 140 format (44x,'/',23x,'\'')
132 150 format (42x,'/',27x,'\'')
133 160 format (40x,'/',30x,'\'')
134 170 format (38x,'/',33x,'\'')
135 180 format (37x,'/',35x,'\'')
136 190 format (36x,'/',37x,'\'')
137 200 format (35x,'/',39x,'\'')
138 210 format (34x,'/',41x,'\'')
139 220 format (33x,'///',6x,'/----\'',7x,'-----',5x,'/----\'',6x,'\\\'')
140 230 format (33x,'///',6x,'1',6x,'1',6x,'1+++1',4x,'1',6x,'1',5x,'\\\'')
141 240 format (9x,'+ z out of paper',7x,'/',7x,'\\----\'',7x,'1+ +1',5x,'\
142      ',7x,'\',12x,'\'')
143 250 format (32x,'///',20x,'1+1+1----- + y
144      direction')
145 260 format (32x,'1',21x,'1+++1',19x,'\'',11x,'/')
146 270 format (32x,'1',21x,'1 1 1',19x,'1')
147 280 format (32x,'1',21x,'1 1 1',19x,'1')
148 290 format (32x,'1',21x,'1 1 1',19x,'1')
149 300 format (32x,'1',21x,'1 1 1',19x,'1')
150 310 format (33x,'\',20x,'1 1 1',19x,'/')
151 320 format (54x,'1 1 1',18x,'/')
152 330 format (54x,'1 1 1')
153 340 format (54x,'1 1 1')
154 350 format (34x,'-----1 1 1-----')
155 360 format (34x,'1 +++++',14x,'1',15x,'+++++ 1')
156 370 format (34x,'1 + 2 +',14x,'1',15x,'+ 3 + 1')
157 380 format (34x,'1 +++++',14x,'1',15x,'+++++ 1')
158 390 format (34x,'1 -----1 -----')
159 400 format (5(56x,'1'),56x,'1')
160 410 format (54x,'1 /')
161 420 format (55x,'1/')
162 430 format (53x,'+ x direction')
163 440 format (1h1,'plexiglass t-plate coordinate system-orientation and
164      ocation of bee bees')
165 c
166      end
```

NUMBER OF ERRORS = 0 NUMBER OF WARNINGS = 0

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

FORTRAN 77/UX HP92430A.07.04 COPYRIGHT HEWLETT-PACKARD CO. 1983.  
Thu Jul 26 14:21:16 1990

```
1      subroutine prism2 (xop,xsp,cp)
2  c
3  c
4  c This subroutine establishes the lab coordinate system and
5  c calculates both the ap and lateral camera orientations and
6  c positions in the lab.
7  c
8  c Inputs are all initial guesses
9  c xo = vector from lab to x-ray origin in x-ray coordinates
10 c
11 c xs = vector from source origin to x-ray origin in x-ray coordinates
12 c
13 c c = transformation matrix from [lab/xray]?
14 c
15 c Outputs are the best estimates of the three inputs above?
16 c
17 c
18 c
19 c Revised December 1987
20 c
21 c The following assumptions were made:
22 c
23 c (1) xs is the same for ap and lateral calibration and is
24 c constant for all runs.
25 c The value of xs: 0.0, 0.0, 150.
26 c
27 c (2) xo is different for ap and lateral calibration but is
28 c constant for all runs.
29 c The values of xo are: (ap) 20., 0.0, 10.; (lat) 0.0, 0.0, 10.
30 c
31 c (3) c is different for ap and lateral calibrations but is
32 c constant for all runs.
33 c The values of c are:
34 c (ap) (lat)
35 c           -1.0, 0.0, 0.0          0.0, 0.0, 1.0
36 c           0.0, 0.0, 1.0          1.0, 0.0, 0.0
37 c           0.0, 1.0, 0.0          0.0, 1.0, 0.0
38 c
39 c*IF ANY OF THESE ASSUMPTIONS ARE WRONG, THIS ROUTINE MUST BE MODIFIED*
40 c
41 c
42 c
43 c     real xop(3,2), xsp(3,2), cp(3,3,2), xpp(2,13,2), pxp(2,2)
44 c     real xt(3,13), a(9,10), xp(2,13), c(3,3), xo(3), xs(3), rt(3)
45 c     real w(10),cc(3,3),xxo(3),xxs(3)
46 c     integer ja(9)
47 c
48 c     data cc /-1.0.0.0,0.0.0.0,0.0.1.0.0.0.0.1.0.0.0/
49 c     data xxo /20.0.0.0,10.0/
50 c     data xxs /0.0.0.0,150.0/
51 c     data xt /5.08,0.,5.08,25.4,0.,5.08,25.4,0.,25.4,5.08,0.,25.4,0.,6.
52 c           20,25.4,0.,5.08,5.08,0.,25.4,5.08,0.,25.4,25.4,5.08,25.4,15.24,15.
53 c           24,15.24,15.24,25.4,5.08,15.24,15.24,15.24,5.08,15.24,15.24,25.4/
54 c
55 c     9 format(3(f7.3,2x))
56 c
57 c     assign a unit number to the input file
58 c     open(10,file='input')
```

## X-Ray Anthropometry Transformation Program

---

```
59      open(11,file='output')
60  c
61      write (11,230)
62      jj=1
63  10  do 12 i=1,3
64      xo(i)=xxo(i)
65      xs(i)=xxs(i)
66      do 12 j=1,3
67      12  c(i,j)=cc(i,j)
68      if(jj.ne.2) go to 15
69      write(11,240)
70      xo(1)=0.0
71      c(1,1)=0.0
72      c(3,1)=1.0
73      c(1,2)=1.0
74      c(3,2)=0.0
75  15  do 20 i=1,13
76      do 20 j=1,2
77  20  xp(j,i)=999.
78  c
79  c  read the required input date
80  c
81      read(10,260) (xp(1,i),i=1,13)
82      read(10,260) (xp(2,i),i=1,13)
83  c
84      do 30 i=1,13
85      xp(1,i)=xp(1,i)*2.54
86      xp(2,i)=xp(2,i)*2.54
87  30  continue
88      nl=0
89  40  err=0.
90      do 50 i=1,9
91      ja(i)=0
92      do 50 j=1,10
93  50  a(i,j)=0.
94      do 100 i=1,13
95      if (xp(1,i).ge.99..or.xp(2,i).ge.99.) go to 100
96      do 60 j=1,3
97      rt(j)=0.
98      do 60 k=1,3
99  60  rt(j)=rt(j)+xt(k,i)*c(j,k)
100     w(1)=xs(3)/(xs(3)-rt(3)-xo(3))
101     w(2)=0.
102     w(3)=w(1)*(rt(1)+xo(1)-xs(1))/(xs(3)-rt(3)-xo(3))
103     w(4)=-w(1)*(rt(3)+xo(3))/xs(3)
104     w(5)=0.
105     w(6)=-w(3)*(rt(3)+xo(3))/xs(3)
106     w(7)=w(3)*rt(2)
107     w(8)=w(1)*rt(3)-w(3)*rt(1)
108     w(9)=-w(1)*rt(2)
109     w(10)=xp(1,i)-xs(1)-w(1)*(rt(1)+xo(1)-xs(1))
110     erp=w(10)**2
111     n=0.
112  70  do 80 k=1,10
113      do 80 j=1,9
114  80  a(j,k)=a(j,k)+w(k)*w(j)
115      err=err+w(10)**2
116      if (n.eq.1) go to 90
117      n=1
118      w(2)=w(1)
119      w(1)=0.
120      w(3)=w(2)*(rt(2)+xo(2)-xs(2))/(xs(3)-rt(3)-xo(3))
```

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

---

```

121      w(5)=w(4)
122      w(4)=0.
123      w(6)=-w(3)*(rt(3)+xo(3))/xs(3)
124      w(7)=w(3)*rt(2)-w(2)*rt(3)
125      w(8)=-w(3)*rt(1)
126      w(9)=w(2)*rt(1)
127      w(10)=xp(2,i)-xs(2)-w(2)*(rt(2)+xo(2)-xs(2))
128      erp=erp+w(10)**2
129      go to 70
130  90  if (nl.eq.20) write (11,270) i,xp(1,i),xp(2,i),erp
131  100  continue
132  call axeb (a,9,10,ja)
133  ao=0.
134  do 110 j=1,3
135  1  ao=ao+a(j+6,10)**2
136  1  xo(j)=xo(j)+a(j,10)
137  1  110  xs(j)=xs(j)+a(j+3,10)
138  ao=sqrt(ao)
139  if (ao.lt..25) go to 130
140  do 120 j=7,9
141  120  a(j,10)=a(j,10)*.25/ao
142  130  r=sqrt(1.-a(9,10)**2)
143  do 140 j=1,3
144  p=c(1,j)
145  c(1,j)=p*r-c(2,j)*a(9,10)
146  140  c(2,j)=p*a(9,10)+c(2,j)*r
147  r=sqrt(1-a(8,10)**2)
148  do 150 j=1,3
149  p=c(1,j)
150  c(1,j)=p*r+c(3,j)*a(8,10)
151  150  c(3,j)=r*c(3,j)-p*a(8,10)
152  r=sqrt(1-a(7,10)**2)
153  do 160 j=1,3
154  p=c(2,j)
155  c(2,j)=p*r-c(3,j)*a(7,10)
156  160  c(3,j)=p*a(7,10)+c(3,j)*r
157  write (11,280) c,xo,xs,err
158  nl=nl+1
159  if (nl.le.20) go to 40
160  do 170 i=1,3
161  xsp(i,jj)=xs(i)
162  xop(i,jj)=xo(i)
163  do 170 j=1,3
164  170  cp(i,j,jj)=c(i,j)
165  do 180 i=1,13
166  xpp(1,i,jj)=xp(1,i)
167  180  xpp(2,i,jj)=xp(2,i)
168  jj=jj+1
169  if (jj.lt.3) go to 10
170  write (11,290)
171  do 210 i=1,13
172  ptp=0.
173  do 190 j=1,2
174  do 190 k=1,2
175  ptp=ptp+xpp(j,i,k)
176  190  ppx(j,k)=xpp(j,i,k)
177  if (ptp.gt.99.) go to 210
178  call pxry2 (w(1),pxp,r,w(4))
179  erp=0.
180  do 200 j=1,3
181  200  erp=erp+(w(j)-xt(j,i))**2
182  write (11,300) i,(xt(j,i),j=1,3),(w(k),k=1,3),erp

```

## *X-Ray Anthropometry Transformation Program*

---

```
183 210 continue
184 c
185 230 format ('1 set up parameters - a-p projection',a6)
186 240 format ('1 set up parameters - lateral projection',a6)
187 250 format (3f10.0)
188 260 format (13(f7.3,1x))
189 270 format (5x,i3,2f10.6,f10.7)
190 280 format (5x,9f10.5)
191 290 format ('1 back-check multiply acquired cal points'///)
192 300 format (5x,i3,6f10.4,f10.7)
193 c
194 end
```

NUMBER OF ERRORS = 0 NUMBER OF WARNINGS = 0

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

FORTRAN 77/UX HP92430A.07.04 COPYRIGHT HEWLETT-PACKARD CO. 1983.  
Thu Jul 26 14:21:30 1990

```
1      subroutine pxry2 (x,a,r,xr)
2  c
3  c subroutine pxryz
4  c
5  c
6  c This subroutine calculates the best least square coordinates
7  c in the lab (x(i),i=1,2,3) consistent with the ap & lateral x-ray
8  c coordinates of all bee bee's and possible right-left ambiguities
9  c arising from the lateral x-ray view.
10 c
11 c input consist of:
12 c   a(2,2) = the ap,x; ap,y; lat,x; lat,y; measured x-ray coordinates
13 c             of the ith bee bee. i=1,7 for neck monut, i=1,13 for head
14 c             mount. Also known a sp(2,2,13) in subroutine rlsrch and
15 c             as spi(4,13) in the main program.
16 c
17 c   xo(3,2) = vectors from lab to x-ray origin (ap:lat) in x-ray coord's
18 c
19 c   xs(3,2) = vector from camera origin to x-ray origin in x-ray coord's
20 c
21 c   c(3,3,2)= transformation matrixces from [lab/x-ray]ap:[lab/xray]lat.
22 c
23 c output consist of:
24 c   x(3) = best least squares estimate of bee bee in lab coordinates
25 c
26 c   r = sum of the square of the error between the measured
27 c       x-ray coordinates and the x-ray coordinates obtained by
28 c       projection of the best least squares position of the bee
29 c       bee onto both x-ray film planes.
30 c
31 c   r = [ap(x,y)measured-ap(x,y)theory]**2.0 +
32 c       [lat(x,y)measured-lat(x,y)theory]
33 c
34 c   xr(2,2) = the x-ray coordinates obtained by projection of the
35 c       calculated position of the bee bee onto the x-ray film
36 c       planes. It is called xtp(4) in the main program.
37 c       It is used in the neck calculations.
38 c
39 c bench marks of intrest consist of:
40 c   x(1) = x(1)+det(2)/det(1) the sum of the change in the x
41 c           component of the lab coordinates of the bee bee for each
42 c           iteration for the best coordinate calculation routine.
43 c
44 c   e = the magnitude of the change squared of x(3) above,
45 c       calculated for each iteration of the best coordinates
46 c       routine.
47 c
48 c
49 c common /blk2/ xo(3,2),xs(3,2),c(3,3,2)
50 c     real x(3), a(2,2), xr(2,2), w(4), gm(3,4), det(4), xx(3)
51 c     nc=0
52 c     do 10 i=1,3
53 10  x(i)=0.
54 20 do 30 i=1,3
55      do 30 j=1,4
56 30  gm(i,j)=0.
57      r=0.
58      nc=nc+1
```

## X-Ray Anthropometry Transformation Program

---

```
59      do 60 i=1,2
60      do 40 j=1,3
61      xx(j)=xo(j,i)-xs(j,i)
62      do 40 k=1,3
63 40  xx(j)=xx(j)+c(j,k,i)*x(k)
64      do 60 j=1,2
65      w(4)=a(j,i)+xx(j)/xx(3)*xs(3,i)-xs(j,i)
66      r=r+w(4)**2
67      do 50 k=1,3
68 50  w(k)=-c(j,k,i)-c(3,k,i)*xx(j)/xx(3)*xs(3,i)/xx(3)
69      do 60 k=1,3
70      do 60 l=1,4
71 60  gm(k,l)=gm(k,l)+w(k)*w(l)
72      do 80 i=1,4
73      det(i)=gm(1,1)*(gm(2,2)*gm(3,3)-gm(2,3)-gm(3,2))+gm(1,2)*(gm(2,3)
74 1)*gm(3,1)-gm(2,1)*gm(3,3))+gm(1,3)*(gm(2,1)*gm(3,2)-gm(3,1)*gm(2,2
75 2))
76      if (i.eq.4) go to 80
77      do 70 k=1,3
78      e=gm(k,i)
79      gm(k,i)=gm(k,4)
80 70  gm(k,4)=e
81 80  continue
82      x(1)=x(1)+det(2)/det(1)
83      x(2)=x(2)-det(3)/det(1)
84      x(3)=x(3)+det(4)/det(1)
85      e=(det(2)**2+det(3)**2+det(4)**2)/det(1)**2
86      if (e.gt..00001.and.nc.le.10) go to 20
87      do 100 i=1,2
88      do 90 j=1,3
89      xx(j)=xo(j,i)-xs(j,i)
90 c
91 c
92 c where xo(j,i) = (lab origin - xray origin) in xray coordinates
93 c xs(j,i) = (xray source - xray origin) in xray coordinates
94 c
95 c
96      do 90 k=1,3
97 90  xx(j)=xx(j)+c(j,k,i)*x(k)
98 c
99 c  xx(j) = (point - source) in xray coordinates
100   do 100 j=1,2
101   c
102 100  xr(j,i)=xs(j,i)-xx(j)/xx(3)*xs(3,i)
103 c
104 c -xx(j)/xx(3)*xs(3,i) = (theoretical xray coordinates of a point
105 c minus source coordinates) in xray system
106 c xr(j,i) = theoretical xray coordinates
107 c
108      return
109 c
110 end
```

NUMBER OF ERRORS = 0 NUMBER OF WARNINGS = 0

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

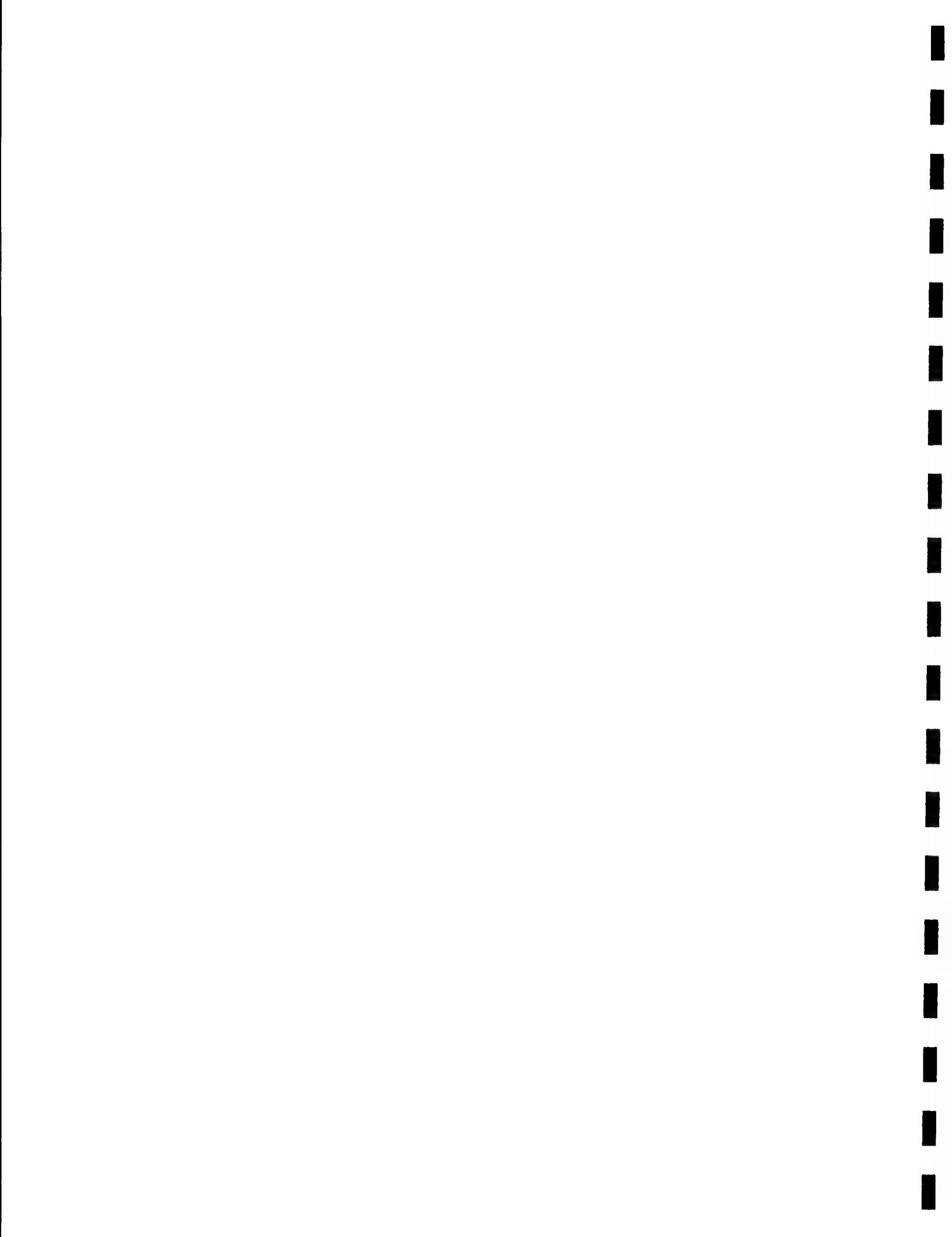
FORTRAN 77/UX HP92430A.07.04 COPYRIGHT HEWLETT-PACKARD CO. 1983.  
Thu Jul 26 14:21:31 1990

```
1      subroutine rlsrch (j,k,sp)
2  c
3  c subroutine rlsrch
4  c
5  c
6  c This subroutine gathers the ap and lat x-ray coordinates of the
7  c anatomical instrument bee bee's. It then generates additional points
8  c by making an alternate switch of the possible ambiguities which
9  c could arise when varying certain bee bee locations on the lateral
10 c x-rays.
11 c
12 c input and output consist of:
13 c sp(2,2,13) = t the first 28 elements of this array contain the
14 c measured coordinates of the ap and lat views of the
15 c the subject's bee bee's, and are passed to this
16 c subroutine from the main program. The last 24 elements
17 c array are empty and will receive the right-left/left-
18 c right switch of coordinates performed by this
19 c subroutine.
20 c
21 c j = the index of the right side bee bee on the lateral
22 c view as obtained from measurement.
23 c
24 c k = the index of the right-side bee bee so labeled by
25 c this subroutine and has identical x-ray coordinates
26 c as the measured labeled left side bb. A least square
27 c solution with accompanying error for both labeled bee
28 c bee's in lab coordinates is determined in subroutine
29 c pxry2. If the proper switch is set in subroutine option
30 c the discrepancies will be automatically corrected there
31 c
32 c
33     real sp(2,2,13)
34     l=k+1
35     do 10 i=1,2
36       sp(i,1,k)=sp(i,1,j)
37       sp(i,2,k)=sp(i,2,j+1)
38       sp(i,1,1)=sp(i,1,j+1)
39   10  sp(i,2,1)=sp(i,2,j)
40       return
41 c
42     end
```

NUMBER OF ERRORS = 0 NUMBER OF WARNINGS = 0

## **APPENDIX B**

**Listing of "IOUPDATE"**



## *X-Ray Anthropometry Transformation Program*

---

```
echo Procedure to update xray input file and to execute main program
cp $1 input
cp output $2
cp update $3
echo Files Saved - Executing Main Xray-Anthropometry Program
mxray
```

## **APPENDIX C**

### **Listing of "INPUT"**

### **X-Ray Anthropometry Transformation Program**

---

H00209	10888	1101	2201	0	0	0	0				
10.386	1.648	1.532	10.273	999.000	999.000	999.000	999.000	9.278	4.890		
1.144	4.973	4.818									
-1.914	-1.944	6.812	7.015	999.000	999.000	999.000	999.000	2.285	2.327		
2.359	-2.526	7.180									
999.000	999.000	999.000	999.000	2.924	2.415	11.212	11.315	11.119	5.850		
-.286	5.800	5.935									
999.000	999.000	999.000	999.000	6.814	-1.942	-2.286	6.697	2.051	1.813		
1.548	-3.216	6.883									
.712	6.362	.994	4.576	2.565	-.581	5.697					
3.036	3.823	4.461	4.814	5.660	2.627	2.389					
1.778	3.033	7.023	7.421	11.146	10.489	10.542					
2.356	3.491	3.935	4.366	5.072	1.787	1.945					
4.729	4.216	5.040	1.992	7.698							
1.597	-1.608	4.446	4.522	4.175							
6.308	11.900	-.370	2.328	3.053							
1.045	-1.989	4.237	4.010	3.804							
2.304	1.575	6.920	1.459								
10.223	-.059	6.750	.369	6.984	.860						

## **APPENDIX D**

### **Listing of "OUTPUT"**

## X-Ray Anthropometry Transformation Program

---

1	set up parameters - a-p projection								
	-.99885	-.01122	-.04663	-.04612	-.04255	.99803	-.01318	.99903	.04199
	31.38578	-9.44438	14.62792	29.17912	.70755	173.346161816.69006			
	-.98938	-.01484	-.14458	-.14440	-.01310	.98943	-.01658	.99980	.01082
	30.64409	-8.72047	19.82321	15.29855	7.45546	185.71568	.97425		
	-.98711	-.01351	-.15944	-.15924	-.01517	.98712	-.01575	.99979	.01283
	29.98223	-8.62734	19.63002	11.52374	7.35874	184.20836	.66993		
	-.98781	-.01232	-.15518	-.15487	-.02303	.98767	-.01574	.99966	.02084
	30.08084	-8.78918	19.42020	12.39414	5.87360	184.01617	.01011		
	-.98702	-.01233	-.16015	-.15984	-.02284	.98688	-.01583	.99966	.02058
	29.97732	-8.78477	19.50641	11.44204	5.90834	184.14708	.00986		
	-.98724	-.01202	-.15878	-.15845	-.02481	.98705	-.01581	.99962	.02259
	30.00607	-8.82524	19.45057	11.70691	5.53493	184.10239	.00981		
	-.98704	-.01204	-.16004	-.15971	-.02468	.98685	-.01583	.99962	.02244
	29.97989	-8.82260	19.47390	11.46572	5.55905	184.13451	.00979		
	-.98710	-.01196	-.15967	-.15933	-.02516	.98690	-.01582	.99961	.02293
	29.98773	-8.83229	19.45997	11.53778	5.46947	184.12390	.00979		
	-.98705	-.01197	-.15998	-.15965	-.02511	.98685	-.01583	.99961	.02287
	29.98128	-8.83130	19.46601	11.47830	5.47863	184.13174	.00979		
	-.98706	-.01195	-.15988	-.15955	-.02522	.98687	-.01583	.99961	.02299
	29.98337	-8.83362	19.46251	11.49755	5.45714	184.12914	.00979		
	-.98705	-.01196	-.15996	-.15962	-.02521	.98686	-.01583	.99961	.02297
	29.98177	-8.83331	19.46407	11.48277	5.46002	184.13101	.00979		
	-.98706	-.01195	-.15993	-.15960	-.02523	.98686	-.01583	.99961	.02300
	29.98231	-8.83387	19.46320	11.48775	5.45484	184.13026	.00979		
	-.98705	-.01195	-.15995	-.15962	-.02523	.98686	-.01583	.99961	.02300
	29.98191	-8.83378	19.46358	11.48410	5.45568	184.13075	.00979		
	-.98705	-.01195	-.15995	-.15961	-.02524	.98686	-.01583	.99961	.02300
	29.98207	-8.83394	19.46332	11.48550	5.45420	184.13048	.00979		
	-.98705	-.01195	-.15995	-.15961	-.02524	.98686	-.01583	.99961	.02300
	29.98194	-8.83392	19.46347	11.48433	5.45441	184.13077	.00979		
	-.98705	-.01195	-.15995	-.15961	-.02524	.98686	-.01583	.99961	.02300
	29.98200	-8.83394	19.46340	11.48492	5.45422	184.13078	.00979		
	-.98705	-.01195	-.15995	-.15961	-.02524	.98686	-.01583	.99961	.02300
	29.98197	-8.83393	19.46343	11.48462	5.45431	184.13074	.00979		
	-.98705	-.01195	-.15995	-.15961	-.02524	.98686	-.01583	.99961	.02300
	29.98199	-8.83392	19.46343	11.48482	5.45435	184.13080	.00979		
	-.98705	-.01195	-.15995	-.15961	-.02524	.98686	-.01583	.99961	.02300
	29.98202	-8.83393	19.46336	11.48503	5.45432	184.13052	.00979		
	-.98705	-.01195	-.15995	-.15961	-.02524	.98686	-.01583	.99961	.02300
	29.98198	-8.83391	19.46344	11.48470	5.45452	184.13078	.00979		
1	26.1380438	-4.861560	.0008375						
2	4.185920	-4.937760	.0010381						
3	3.891280	17.302479	.0026324						
4	26.093418	17.818100	.0012070						
9	23.566120	5.803900	.0008343						
10	12.420600	5.910580	.0001535						
11	2.905760	5.991860	.0020173						
12	12.631420	-6.416040	.0005440						
13	12.237720	18.237200	.0005259						
	-.98705	-.01195	-.15995	-.15961	-.02524	.98686	-.01583	.99961	.02300
	29.98200	-8.83391	19.46340	11.48492	5.45443	184.13060	.00979		
1	set up parameters - lateral projection								
	.04389	-.00293	.99903	.99892	-.01517	-.04393	.01528	.99988	.00227
	3.16778	-8.97930	15.70602	44.45662	16.28766	150.192081487.97229			
	-.08163	-.04970	.99542	.99648	-.02306	.08056	.01895	.99850	.05141
	2.51471	-8.81139	13.89865	23.52338	8.10413	149.23682	45.38939		
	-.17244	-.05655	.98340	.98496	-.02059	.17153	.01055	.99819	.05925
	1.10893	-8.95892	10.63221	9.24914	6.91589	139.99602	.19440		
	-.15521	-.01355	.98779	.98780	-.01486	.15501	.01258	.99980	.01570
	1.51912	-8.40127	11.46772	12.50575	13.29204	140.83527	.17037		
	-.17620	-.01257	.98427	.98429	-.01364	.17603	.01121	.99983	.01478
	1.27281	-8.39783	10.80466	9.45897	13.41144	138.94174	.04211		
	-.17219	-.00647	.98504	.98500	-.01288	.17210	.01158	.99990	.00859
	1.32988	-8.31788	10.97785	10.08479	14.31426	139.26227	.03585		
	-.17428	-.00675	.98467	.98463	-.01284	.17419	.01147	.99989	.00889
	1.30591	-8.32315	10.91266	9.78224	14.26912	139.08342	.03520		
	-.17391	-.00628	.98474	.98469	-.01279	.17382	.01150	.99990	.00841
	1.31041	-8.31674	10.92883	9.83791	14.33949	139.11691	.03517		
	-.17407	-.00631	.98471	.98467	-.01279	.17398	.01149	.99990	.00844
	1.30856	-8.31732	10.92386	9.81488	14.33423	139.10335	.03517		
	-.17404	-.00628	.98472	.98467	-.01278	.17395	.01150	.99990	.00841
	1.30892	-8.31686	10.92514	9.81932	14.33915	139.10625	.03517		
	-.17405	-.00628	.98472	.98467	-.01278	.17396	.01149	.99990	.00841

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

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1.30879	-8.31692	10.92477	9.81772	14.33866	139.10519	.03517			
-.17405	-.00628	.98472	.98467	-.01278	.17396	.01149	.99990	.00841	
1.30881	-8.31689	10.92485	9.81797	14.33892	139.10538	.03517			
-.17405	-.00628	.98472	.98467	-.01278	.17396	.01149	.99990	.00841	
1.30880	-8.31689	10.92483	9.81786	14.33896	139.10530	.03517			
-.17405	-.00628	.98472	.98467	-.01278	.17396	.01149	.99990	.00841	
1.30882	-8.31689	10.92485	9.81804	14.33901	139.10542	.03517			
-.17405	-.00628	.98472	.98467	-.01278	.17396	.01149	.99990	.00841	
1.30881	-8.31688	10.92487	9.81794	14.33908	139.10548	.03517			
-.17405	-.00628	.98472	.98467	-.01278	.17396	.01149	.99990	.00841	
1.30880	-8.31689	10.92482	9.81780	14.33897	139.10538	.03517			
-.17405	-.00628	.98472	.98467	-.01278	.17396	.01149	.99990	.00841	
1.30880	-8.31688	10.92483	9.81788	14.33910	139.10533	.03517			
-.17405	-.00628	.98472	.98467	-.01278	.17396	.01149	.99990	.00841	
1.30879	-8.31687	10.92483	9.81777	14.33912	139.10532	.03517			
-.17405	-.00628	.98472	.98467	-.01278	.17396	.01149	.99990	.00841	
1.30880	-8.31687	10.92483	9.81783	14.33912	139.10533	.03517			
-.17405	-.00628	.98472	.98467	-.01278	.17396	.01149	.99990	.00841	
1.30880	-8.31688	10.92485	9.81791	14.33909	139.10535	.03517			
5	7.426960	17.307560	.0081797						
6	6.134100	-4.932680	.0075463						
7	28.478479	-5.806440	.0020279						
8	28.740099	17.010380	.0026793						
9	28.242260	5.209540	.0089725						
10	14.858999	4.605020	.0010488						
11	-.726440	3.931920	.0001521						
12	14.732000	-8.168640	.0001277						
13	15.074900	17.482819	.0044327						
	-.17405	-.00628	.98472	.98467	-.01278	.17396	.01149	.99990	.00841
1	1.30881	-8.31688	10.92484	9.81791	14.33903	139.10544	.03517		

1 back-check multiply acquired cal points

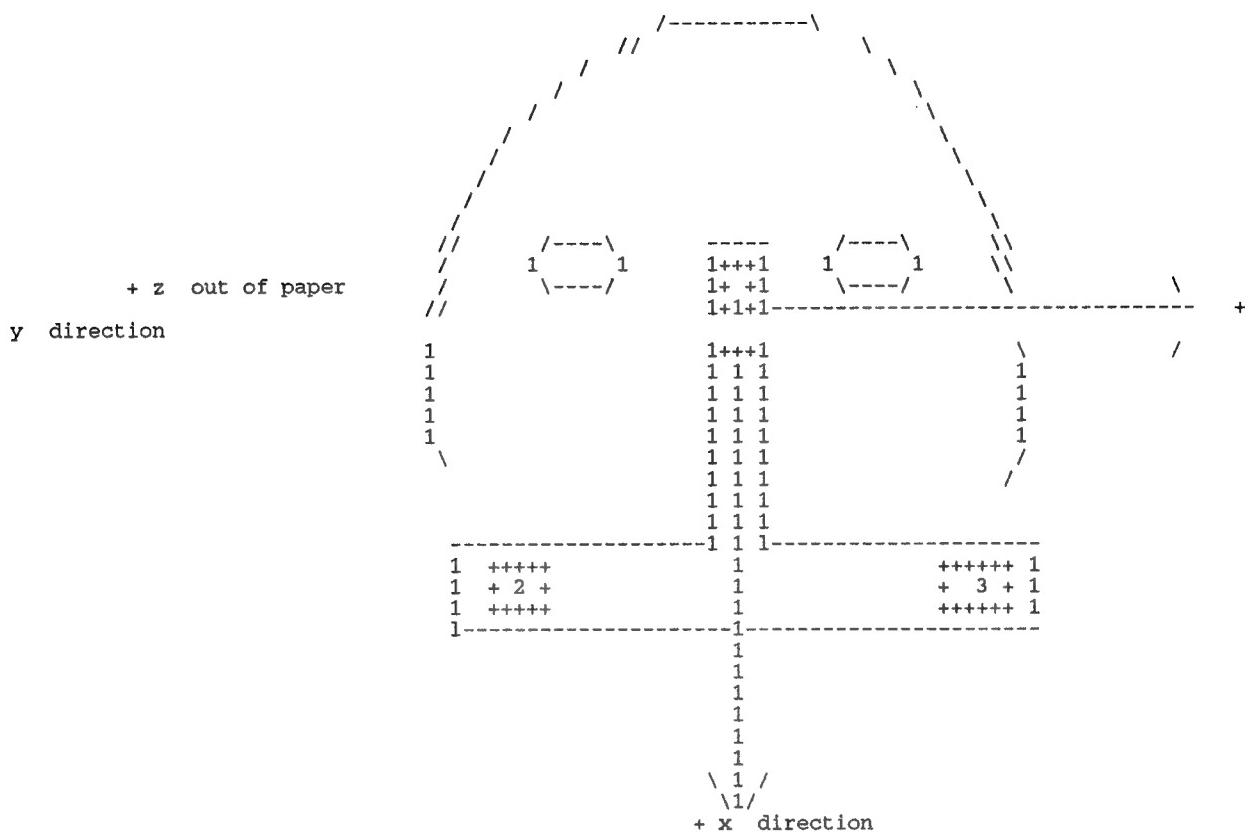
9	5.0800	25.4000	15.2400	5.0860	25.3208	15.2453	.0063406
10	15.2400	15.2400	15.2400	15.2532	15.2224	15.2299	.0005861
11	25.4000	5.0800	15.2400	25.3830	5.0853	15.2235	.0005880
12	15.2400	15.2400	5.0800	15.2349	15.2394	5.0657	.0002298
13	15.2400	15.2400	25.4000	15.2366	15.2699	25.4122	.0010533

1 head anatomical to tee-plate transformation  
subject H00209 mount 1101 date APR88 notes HEAD DATA

measured x-ray coordinates				lab coordinates				sum of	
the squares of the	i	i	i	i	i	i	i		
difference between the	i	i	i	i	i	i	i		
measured and best estimate	i	i	i	i	i	i	i		
x-ray coordinates	i	ap	lat	i	i	i	of		
	i		i	i	i	i	i		
0	i	x	y	x	y	z	e		
1	r.audit	1.8085	7.7114	4.5161	5.9842	25.4972	9.0287	16.8190	.0031
2	l.audit	16.1595	9.7104	7.7038	8.8671	12.9452	8.9599	18.3497	.0029
3	r.orbit	2.5248	11.3309	17.8384	9.9949	22.8189	18.4356	19.8274	.0008
4	l.orbit	11.6230	12.2276	18.8493	11.0896	15.3324	18.3475	20.4165	.0003
5	c.t-plt	6.5151	14.3764	28.3108	12.8829	18.0769	25.8508	22.0095	.0007
6	r.t-plt	-1.4757	6.6726	26.6421	4.5390	24.6725	25.0227	16.1485	.0015
7	l.t-plt	14.4704	6.0681	26.7767	4.9403	12.2026	24.4568	15.5134	.0011
0	right/left switch follow below								
8	r.audit	1.8085	7.7114	7.7038	8.8671	24.9421	11.2660	18.0525	3.7105
9	l.audit	16.1595	9.7104	4.5161	5.9842	13.3660	6.4079	17.0826	3.8006
10	r.orbit	2.5248	11.3309	18.8493	11.0896	22.6515	19.1498	20.2684	.6052
11	l.orbit	11.6230	12.2276	17.8384	9.9949	15.4720	17.5776	19.9732	.5842
12	r.t-plt	-1.4757	6.6726	26.7767	4.9403	24.6381	25.1154	16.3048	.0536
13	l.t-plt	14.4704	6.0681	26.6421	4.5390	12.2300	24.3515	15.3570	.1033
	option 1 vs 8 not indicated								
	option 3 vs 10 not indicated								
	option 6 vs 12 not indicated								

1plexiglass t-plate coordinate system-orientation and location of bee bees

## X-Ray Anthropometry Transformation Program



	x	y	z
bb 1	.0000	.0000	.0000
location of bee bees in instrumentation:	bb 2	6.2860	-6.2860
	bb 3	6.2860	6.2860

1 head anatomical to tee-plate transformation  
 subject H00209 mount 1101 date APR88  
 notes HEAD DATA

anatomy vs lab				instrumentation vs lab			
19.2212	-.01495874	.96531314	.26066601	18.0748	.05787801	-.99769205	-.03550632
8.9943	-.99450266	-.04138450	.09618630	25.8505	-.17667633	-.04524093	.98322874
17.5844	.10363743	-.25779423	.96062553	22.0081	-.98256588	-.05063419	-.17888699

instrumentation vs anatomy			
17.4417	-.42753530	-.04194606	.90302497
.8681	-.14475755	.98920941	-.02258578
-.2147	-.89233339	-.14037588	-.42899388
1 t-1	anatomical to tee-plate transformation		
subject H00209	mount 2201	date APR88	
notes NECK DATA			
0bee bee id i	measured x-ray coordinates	i	lab coordinates
the squares of the	i	i	i
			sum of
			i

## NAVAL BIODYNAMICS LABORATORY SOFTWARE DOCUMENTATION

---

difference between the  
 measured and best estimate  
 x-ray coordinates

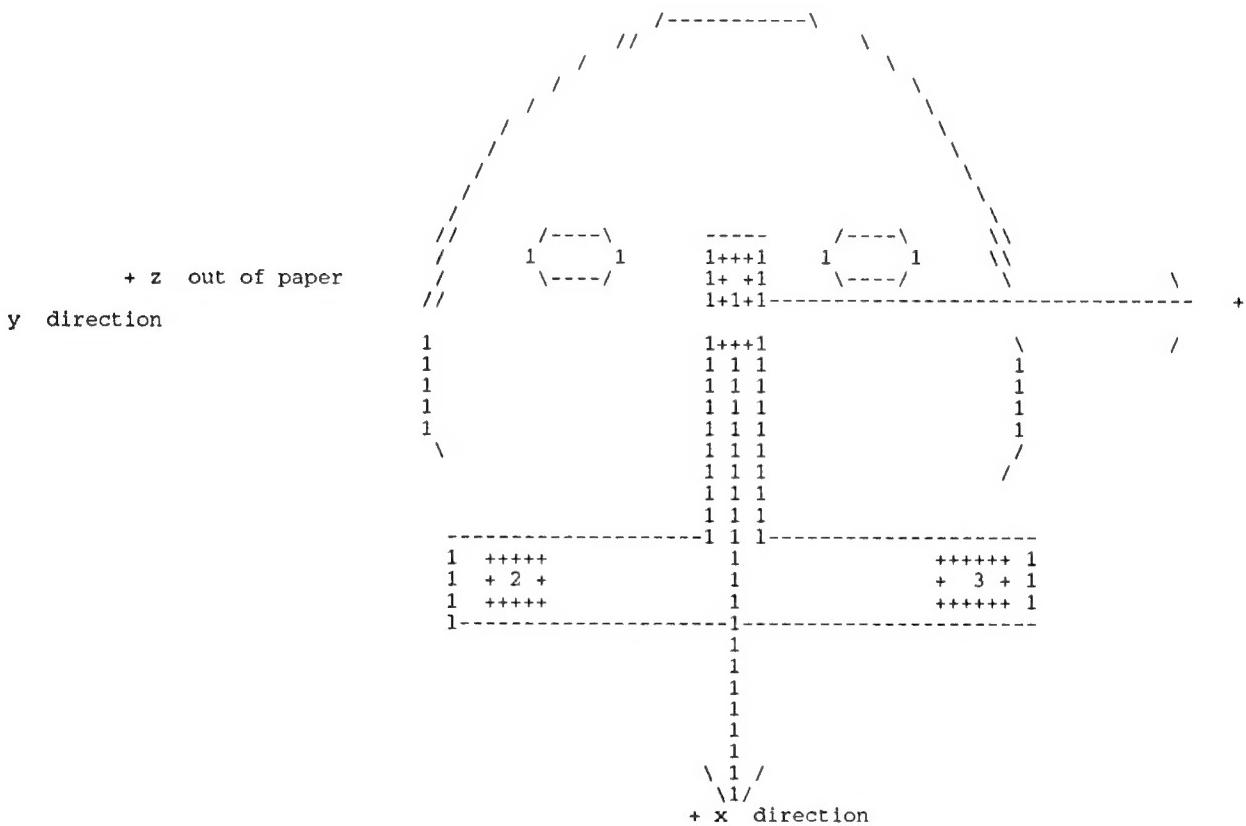
	i	i	i	i	i	i	i
	ap	lat	i	i	x	y	z
0 1 p-spine	12.0117	4.0564	16.0223	2.6543	15.4639	16.1980	13.7306
2 s-notch	10.7086	-4.0843	30.2260	-5.0521	14.7823	27.3728	7.9246
3 c.t-plt	12.8016	11.2928	- .9398	10.7620	16.7990	2.6589	19.8672
4 r.t-plt	5.0597	11.4859	5.9131	10.1854	22.5042	9.4179	19.8886
5 l.t-plt	19.5529	10.6045	7.7546	9.6622	10.0794	8.4461	18.9701
0 6 r.t-plt	5.0597	11.4859	7.7546	9.6622	22.2442	10.7782	19.6704
7 l.t-plt	19.5529	10.6045	5.9131	10.1854	10.2519	6.8995	19.1873
option 4 vs 6 not indicated							
plexiglass t-plate coordinate system-orientation and location of bee bees							

right/left switch follow below

0 6 r.t-plt 5.0597 11.4859 7.7546 9.6622 22.2442 10.7782 19.6704 .2348

7 l.t-plt 19.5529 10.6045 5.9131 10.1854 10.2519 6.8995 19.1873 .0556

1plexiglass t-plate coordinate system-orientation and location of bee bees



		x	y	z	
	bb 1	.0000	.0000	.0000	
	location of bee bees in instrumentation:	bb 2	6.2860	-6.2860	.0000
		bb 3	6.2860	6.2860	.0000
0	articular facets	5.8522	4.0005	17.5768	
0	lateral projection	25.9664	-14.99	17.4422	
		14.9090	24.0173	11.6953	
		15.4044	17.3306	12.8952	
1	t-1 anatomical to tee-plate transformation				
	subject H00209 mount 2201 date APR88				
	notes NECK DATA				

## X-Ray Anthropometry Transformation Program

### anatomy vs lab

14.9090	-.07271960	.98167169	-.17616072
24.0173	-.99600524	-.08065766	-.03831476
11.6953	-.05182123	.17267077	.98361552

### instrumentation vs lab

16.7738	-.07466944	-.99411541	-.07847808
2.6720	.99481934	-.07970386	.06310345
19.8644	-.06898712	-.07335965	.99491668

### instrumentation vs anatomy

-22.5287	.99416876	.00697174	-.10761147
-.4487	-.00322542	.99938363	.03495481
4.2529	.10778888	-.03440393	.99357843

## **APPENDIX E**

### **Listing of "UPDATE"**

**X-Ray Anthropometry Transformation Program**

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H00209	1101	APR88	HEAD DATA	0	0	0	0
17.441742	.868058	-.214688					
-.42753530	-.14475755	-.89233339					
-.04194606	.98920941	-.14037588					
.90302497	-.02258578	-.42899388					
H00209	2201	APR88	BLOCK NECK DATA	0	0	0	0
-22.528715	-.448652	4.252915					
.99416876	-.00322542	.10778888					
.00697174	.99938363	-.03440393					
-.10761147	.03495481	.99357843					